

**Amendments to the Drawings**

Replacement Figure 3B is being transmitted herewith.

**REMARKS**

**Amendment to the Drawings**

The amendment is indicated below and a replacement drawing and annotated sheet for FIG. 3B are submitted herewith.

FIG. 3B has been amended to correct a typographical error. The fifth amino acid from the C-terminus in the Figure is "C" whilst it should be "Q" to correspond with the Sequence List and to correctly translate the nucleic acid sequence listed immediately above the amino acid abbreviation.

The Notice to Comply requested a computer readable from (CRF) copy of the Sequence Listing, a paper copy of the Sequence Listing, as well as an amendment specifically directing its entry into the application, and a statement that the content of the paper and computer readable copies are the same and include no new matter. A copy of the Notice is included with this Amendment.

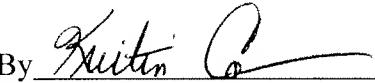
Applicants have identified the error as being an error in the Replacement Drawings that were submitted on June 5, 2009, and not an error in the Sequence Listing. In response to the Notice to Comply, Applicants are submitting this Amendment rather than a Substitute Sequence Listing.

**CONCLUSION**

In view of the above amendments and remarks, it is believed that the application is in condition for allowance and all formal matters have been resolved, and it is respectfully requested that the application be passed to issue. If the Examiner feels that a telephone conference would expedite issuance of this case, the Examiner is invited to call the undersigned.

Respectfully submitted,

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# Annotated Sheet

## FIG. 3B

CAGGGTGACAGAGCCGCTGGGGCTGCCGACCTGGCAACCCCTGTCCCTGCCAGGCCACGATGAGGC  
 R V T E R W R L P T L A T P V I P A L Y **C** Q A T M R 640

TGCCCTGGCTTGAGGCTCAGCCACCGCCAGGCCATCCGGTGCACGGCCGACCTCCGGAGCCCCGACACGACC  
 L P G L E L S H R Q A I P V I H G P T S R E P P D T T 720

TCCCCGGAAACCCCCGGCCGGCACCTCCCCGGAGACCACCCCCCAGCAGGGCTCCACACACAGCCCCCAGGAGCCCCGGGCTC  
 S P D P R A T S P E T T P Q C G S T R S P R S P G S 800

TACCAAGACTTGGCCGCCCTGAGATCTCCCAGGGCTGGCCACGGCAGGGAGAAGTGATCCCACAGGCTCGTCCAAACCTA  
 T R T C R P E I S Q A G P T Q G F V I P T G S S K P 880

CGGGTGACCAAGCTGCCGGCTCTGGACCAAGCAGTGGGACTGGGGCTGGCTGGCCACCTACCAAC  
 T G D Q L P A L W T S S A V L G L L A I P T Y H 960

CTCTGGAAACGTTGCCGGCACCTGGCTGAGGACCTGGGGCTGGCTGGCTTGGCCACCTACCAAC  
 L W K R C R H L A E D G A H P P A S L S S ~~Q~~ P F P L . 1040

AAGGGAAAATAGGTTGGACCCCTTCAAAGCTGAGAACTGGTCGGGGCAACACCTGCCTCCATTCTATTCAAAGTCATCGCT 1120

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